

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Ruddy, David A.
 Wolff, Roger K.
 - (ii) TITLE OF INVENTION: POLYMORPHISMS IN THE REGION OF THE HUMAN HEMOCHROMATOSIS GENE
 - (iii) NUMBER OF SEQUENCES: 24
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
 - (B) STREET: Two Embarcadero Center, 8th Floor
 - (C) CITY: San Francisco
 - (D) STATE: CA
 - (E) COUNTRY: USA
 - (F) ZIP: 94111
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/852,495
 - (B) FILING DATE: 07-MAY-1997
 - (C) CLASSIFICATION: 435
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/724,394
 - (B) FILING DATE: 01-OCT-1996
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/652,265
 - (B) FILING DATE: 23-MAY-1996
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/630,912
 - (B) FILING DATE: 04-APR-1996
 - (C) CLASSIFICATION:
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 235033 base pairs
 - (B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CACACACACA	CACACACACA	CACACACACA	CACACAAATG	AGGTATATAA	AGGGTCTCCT	60
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CAGAAGAAAA	ATCAAACAGG	CATTTCAGAG	ATTGAGGCCA	AGAAGTTAAA	TGTCTTAAAT	240
GGGCAGAGCT	TAGCTGCTTG	ATGTGAAAAG	AGACCAGCGT	GGCTGGAACA	GCAAAGGAGA	300
ACAGCAGAAG	AGGTGAACAG	AGGCCAGAGA	TGGTCACTGA	GTGGGCCCTT	AAGTCATGGT	360
AAGGAGTATG	GAGAATGAAT	TATTGCATGT	ATTGAATATG	TAGGTGACGT	GACTCACAGA	420
TACTTTGGAT	TTGTAGAGAT	GAAGGAAATG	TAGCAAGTGA	CACTCTTAGA	ATGTTGATTT	480
GAGTAAATGG	TAGTGTCAGT	TATTGAACTG	GGGAGAACTG	GAAGGGATAA	CAGGCTTAAG	540
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GCTGGGCATG	GTGATGTGCT	TCTGTGGTCC	CAGCTACTCG	GGAGGCTGAG	GCTGAAGAAT	1560
CGCTTAAACC	CGGGGGGTGG	AGGTTGCAGT	GACCCAAGAT	CACACCACTG	CACTCCAGCC	1620
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ATGTTGCCTA	GGCTGGTCTC	TAACTCCTGA	GCTCAAGTGA	TCTGCCCTCC	TCAGTCTCCC	2340
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ACCTTCCTTC	TTTCCTATTT	CTCCATTTTC	TAATGGGAAT	GTCTATTATG	CCTGTTTCAC	3360
CATTGTACCT	TAGAAGCATG	TAACATTTCT	GGTTTCACAC	GTTCAAAGCT	GGAAAGGAAT	3420
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CATGTCTCAG	CCTCCAGAGT	AGCTGGGACT	ACAGGCGCCC	GCCACCACGC	CTGGCTAATT	3720
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CATGCCCGGC	TGGGATGGAA	TAAATTTATC	TTGTATGGGA	GAAGGACATA	CATTTTGGCA	3900
GGTCAAGGAC	AGAATGTTAT	GGACTAAACT	GTGTCCCCCA	AAATTCATTT	ATTAAAACCC	3960
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TTCCAGGCTC	CAAAACTGTG	AGAAAATGAA	TTTCTGTTCC	AAGCCTCTTA	GATATGGAAA	4260
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GCTAAGACAA	TGAAGGATGT	GGTAAAACTT	TACGTCCCAA	CCACATACCA	AAGAGGCTGG	4380
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CATCCAATGA	AGTTCTGACA	TTTCTTCAAC	ATGAGTACAG	TAATTCAATG	CCAGAGAATT	4560

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TGGGAGACCT	CTAGGTTTAG	CATCCTCATC	CACTCGCCCC	AATTTAAATA	GTCCTCCCCA	5640
GGGCCATTCA	GGCAAGGGAG	ATGAAAACTT	GCTCAAGAGT	TGGAATCCAA	CTGAAGCTAC	5700
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CGAGGAATGT	CCTTTGCTTA	GGGACTAGGC	TCTTAGCAGT	ACCTCTTAGG	TAAGAACTGG	5880
TTAACTGGCA	CCTTCTGTGT	TTCTCTGAAG	CTCCCTTTGC	TTAGGGACTA	GGCTCTTAGC	5940
AGTACCTCTT	AGGTAAGAAC	TGGTTAACTG	ACACCTTCTA	TGTGTCTGAA	GCTCCCAGAA	6000
CAAACTGCCA	GTGAAATTTG	GATTTTTGGA	ATATAGTTTC	TTTTTTTTTT	TTACTTTTTG	6060
TTTTGTTGTT	TTTTTTTGAG	AGTCTCACTC	TCACTGCAAC	CTCCCCCTCC	TATATTCAAG	6120
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GGTGAACACC	GCCACACCTG	ACTAATTTGT	GTAGTTTTAT	TAGAGATGGG	GTTTCGCCAT	6420
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CGGGGGACTA	GTCGGAGGAC	CAAACAAGGT	TACCAACACG	TTAGAGTTTT	GCCTTCAATT	9240
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TGCCCCGGTA	ATAGTCTTTT	AACCTAATAT	GCGTCAGTTT	TGATAACAAC	ACTAAGGCAG	11340
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CTCCGCCTCC	CGGGTTCAAG	CCATTCTCCT	GCCTCAGCCT	CCGGAGTAGC	TGGGATTACA	12300
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GTCTTGGAAA	CAGGTTGTCT	ATCCCTGGAC	CCTTTGAGTT	TTCTGTTCAC	TTTCCTTTGG	12840
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CAACCCGCCC	AGTCTTGTTA	GACATGGGGT	CTGTAGTTTC	TAGTAGGTTC	TTGAGTCTAG	25260
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CTGAGGCAAG AGAATTGCTT GAACCCAGGA GGCGGAGGTT GCAGTGAGCC GAAATCGCGC 137460 CACTGCACTC CAGCCTGGGT AACAGAGCAA GGCTCTGTTT CAAAAATAAA TAAATACATA 137520 AATAAATATT TTTTAAAAAA AGAACATCAC TATGCACCCC ATATATACAT ATAATTATTA 137580 TGTCAATTTG AAACATAATT TTGAAAAATG AAAAAATGAA ACACAAATAT GAATCAATCC 137640 TCTCCAAGTT GATATACTTA AAAGGAAAAA AGTCCGAGGG CTTAAACTAT TCAATCAAAA 137700 TTTTATTAAA ATGCTATAGT AATCTGGAAA GTATTTCAGA ATGAATTGGT ATAAGGTTAG 137760 ACACAAAGAT CAGTGAAACA AAACAGAGAA CCCAGAAATA GATTCACACA TCTATGGACA 137820 ACTGGTTTTG ACAAAGGTGT CAAGGCTATT TAATAAGTAA AAAAATCGTC TTTTCAGTAA 137880 ATGTTTCTTG AACAAGTAGA CATCCGGTGT GGGGGAGAGG AGCAGGAGCC TTACCTCAAA 137940 CTTTATGCAA AAATTAACTC AAAATAGACC ATAGACTTAA ATGTAAAAGC TAAAATTATA 138000 AAACTTCTTT AAAAAATAGG AGAAAATCAT CAACACCCTA GGATTAGCAA AGATTTCTTT 138060 AAAACAAAAC AACAGGTTTA TAGTTTATAA AACATAAATA ACAAAATGAT AAATTTCATC 138120 AAAAGTGAAA ATTTGCTTTT CAAAAAACAT TATAAAATGA AAAGCAGGAG GCTGAGGCAT 138180 GAGAATCACT GGAACCCGGG AGCTACAGGT TGCAGTGAGC CAAGATGGTG CCACTGCACT 138240 AAAAGAAAAA GAAAAATCAC AGGCTGAGAG AAAATATTTA TAATACATGT ATCTGACAAA 138360 GGACTCGCAC CTGGAAAATA TAAGGAACCT TATAACTTAG TAAGATGACA AGCCAAAACA 138420 AAGAGTAAAA GTTTTCAACA GACATTTCAC AAAAGAAAAC ATACAAATGG CCAGTATGCA 138480 CATGAAAAGA TTTTAAACAT CATTAGTTAC TAGGGAAATG CAAGTCAAAA CCACAATGAG 138540 ATACTTCACA TTCAACAGAA TAGCTAATGT TAAAAGGACT GACAATCCCC AGGGTGAGCA 138600 AGGGTGTGGA GGAAACTACT CTCATATATT GTGAATGTAA GAGGACAATG TTACAACTAC 138660 TTTGAAAAA GTTTGGCTGT TTCTAACATA AAATTAAACA CTTATACAGC CCAGCAATAT 138720 TTCTGGGTCA TTTCTCCCAG ATAAATGAAC ACATGTCCAT ACTATGACAT GTACAAATGT 138780 TCATACTGGC TTTGTTTCAC AATGCTATAA ACTGGAAACA ACCCACGTGT CCATCAACAG 138840 GTGAATGGGT AAATAAATTG TAATATATCG GCCAGACGCA GTGGTTCATG CCTGTAATCC 138900 CAGAACTTTG GGAGGCCAAG ATGTACGGAT CACCTGAGAT CAGGAGTTTG AGACCAGCCC 138960 ATCCAACATG GTGAAACCCC ATCTCTACTA AAAAATTAGC TGGGCATGGT CACGGGCGCC 139020

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GATTGTTGGT CATCTATTCG ATGTCCCTTC TCTGTTTTTT AATATGAGAA CTCCTGATTT 147180 TCAGCCAACT ACCCTGGAAA AAAAGCTAAT CTTTCTGACT TCTTAAGTGT GGCCATGTAC 147240 TAAATTCTGG CTAATGCAAG GCAAGCCAAA GGTTTTATGA TAGGTTTTAG GACACTAGAG 147300 TAAAAGAGAG CTGTTGCACA CATGCTCTTC ACCCTACTTT TGTGTCCTTT TTTCCATCCT 147360 ACAACTTGGG TTGTGAGTAT GATGGCTGGA ACTTTAGTGG CTCTCTTGGA TCCCAGGGGT 147420 AATTGAGGGG TGGCTGGAAG GAATCTGTGA TTTTCTGGAG TTTCCATACA CAAACAAGAC 147480 CTGGATTTC TGGGCTTCCC AGACTTCCAC ATCTAGACTT GCTTTAAATG GGAGATAAAT 147540 AAACTTGTTT CAGCCACTGT CATTTTGGGC TATTTTATAG AACTTAATCT AATCTTCAAG 147600 GGTACATGAA TTGCTTTTCC TTAAAAAAAA AATCAGCCAT AAAATCATCT TCTTTTTTCT 147660 TTTGTTCCCC ACATTATTTA GTTGGAGCTC TGTAACTTTT TTTTTTTTT TTTTTGAGAC 147720 AAGGTCTTGC TCTGTCACTT AGGCTGGAAT TCAGTGGCAT GACCATGGCT CACTGCAGCC 147780 TTGCCCTCCT AGGCTCAAGC AATCCTCGTC TCAGCCTCCT GAGTAGCTGA AACTAAGGCA 147840 CATGCCACCA TGCCCAGCTA ATTTCTTTC TTTTAGAGAT GGGAGCCTTG CCCAGGCTAG 147900 TCTCAAACTC CTAGCCTCAA GTGATCCTCC CATCTCAGCC TCCCAAAGTG ACAGGATTAC 147960 AGGTGTGAGC CACCATGCCT GGCTGCTCTG TAAGTGTCTG AATTTCATTT TGTATTTATC 148020 AGTCTGTTTA GATTTTCTTT CCCTTCTTGG GTCAGTTAGG CCATTGGTTT CTTTTTAAAG 148080 GTTTTCAAAT TTATTTGCAT CTAATTCTTC AAATTACTCT CAAAATTATT CCAGTATATA 148140 TTCTTTTGTT CCTATTTCT TCTGTATTCT TTATTAAAAT AGCTAATGAT TTATCTAGCA 148200 GGACTTATAT TCTTTCCATA ACTTTCCTGC ACCCCAATTA ATCTCCAATT TTATATTTCT 148260 TCTGGCCTTC CTTATAGTTT CCACAGGTTT ATTTTATTCA TTTTTTAAAA CTTTTATTTA 148320 ATTGTTTATT TTATTATCAT TCTTTCTTAT TCAGCAATCT AAGTGCTTAG GGATATAGAA 148380 TTTCCTCTAA GCAGCATATG CTAGGCTTTA ACAATGTTAG GGAGGCCTCC CCTTTCTGGG 148440 GAAGACCACA CTTACATTAA CACAGGACTG TGGGATGCCA AGAGGTAGAG AAGAGCTTAT 148500 GAATATCCAG ATTACATCTT CACTGATCCT GCACAAAGGT GGGGTTCCTC GGTTACCCAC 148560 TGGGTCCTAT TACCCAAGTC TGGGTCAGCA TACCGAGACT ACGGGTATAT AGAACAAGTG 148620 CAACTGGCGA TAATCCTTCT GTTGGGGAGA AAAATCTTTT TTTTCTATTC ATCTTAGGTT 148680 CTCCATCTGT GGCCCTATCA AGTAGACTAA CAAAAGACAG ATTGACAAGA CAGAAACAAA 148740

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Chhumanaasa	CTC A A CCC A C	3 3 mmm.cc 3 cc	COACCCCCCC	CCCC A mm A C *	COOR OF COOR	104160
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GAG	BACGGAGT	TTTGTTGTCA	TTGCCCAGGC	TTAGTGCAGT	AGCGCGATCT	CACCTCACTG	198840
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TGACTGTTTA	GCCATTTTAG	AGTGTGGCAT	ACGTGGCTGC	TGGCATACAG	AGGTTGGATG	229140
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GCTCTTGAAC	TGGGATTTAC	ATCATCAGTT	CCTCTGGTTC	TCAGGCCTTC	AGATTCAGAC	234240
TGAATCATAC	CACCAGCTTT	CCTGGGTCTC	CAGCTTGCAG	ATTACAGATC	ATGGGACTCC	234300
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GTGTGTGTGT GTG

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 237326 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

CACACACAC CACACACACA CACACACACA CACAAATGAG GTATATAAAG GGTCTCCTAA 60 120 AATGTCATCT GATATTTGTT ATTTCATATT CTCAGATTTT TAATCCATTT AGGTAGGTCT ATTTTAGATA GCCTTGTCTG AAACAGAGCT GGGACCTGAT GAGTGAAAAT GAGCTCACCA 180 GAAGAAAAT CAAACAGGCA TTTCAGAGAT TGAGGCCAAG AAGTTAAATG TCTTAAATGG 240 GCAGAGCTTA GCTGCTTGAT GTGAAAAGAG ACCAGCGTGG CTGGAACAGC AAAGGAGAAC 300 AGCAGAAGAG GTGAACAGAG GCCAGAGATG GTCACTGAGT GGGCCCTTAA GTCATGGTAA 360 GGAGTATGGA GAATGAATTA TTGCATGTAT TGAATATGTA GGTGACGTGA CTCACAGATA 420 CTTTGGATTT GTAGAGATGA AGGAAATGTA GCAAGTGACA CTCTTAGAAT GTTGATTTGA 480 GTAAATGGTA GTGTCAGTTA TTGAACTGGG GAGAACTGGA AGGGATAACA GGCTTAAGGA 540 GCACGTTTAT TCCTGTGTCT TGGAAGTGTT TAGGGTGAAA GACCTATTAG AGTTCTAAAT 600 GGAGATGTCA AGTGAAAATG TGGCTACACA CATTTGCATT TCAGAAAAAA GGTCAGGCTG 660

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1080	TCCAAGCTGA	GTTATGGCAG	GTAGTATTTT	CCACCAGTCT	ATCGTTTAAG	ATAAATGTCT
1140	ATGGGGAAGT	ATACCTAAAC	GCAACAACAA	GTGGGATGCT	GTACCCAGGC	CTAAGTTTTG
1200	CTAGAAAAAG	GAGGTTCATA	GGAAGAGTTT	GGTAAAGGCT	ATTGGTGATG	GGCTTTGGAA
1260	GGCAAAGGCT	AGGCAATTCT	TGGACATTAA	GAAAGAAATA	AGGGACTATT	CCAATTGTGA
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1380	GGCGTGGTGG	CTTTAGGCCA	AAAATATGGA	TATGCTGGTT	GAATATTAAA	ATCATGGATA
1440	GGTCGGGAGT	CAGATCACGA	GCTGAGGGCA	ACTTTGGGAG	TAATCTCAGC	CTCACGCCTG
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1560	TGAAGAATCG	AGGCTGAGGC	GCTACTCGGG	TGTGGTCCCA	GATGTGCTTC	TGGGCATGGT
1620	CTCCAGCCTG	CACCACTGCA	CCCAAGATCA	GTTGCAGTGA	GGGGGTGGAG	CTTAAACCCG
1680	ATATATATGG	CAAAAAATAT	CACACACA	CTCCCCCGC	CAGGACTCCA	GGATACAGAG
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1920	GAGAGGAAAG	AGTGAAATGT	TACTGCTTAA	AGGTCCTCCT	GGTGTGATTT	AAGTATTGAA
1980	ATTTGGGATA	GAACTTGGAG	AACACAATCA	TTTTAAGCAA	AAGAAGGAAT	AGCCGAAATA
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2220	GACTACATGT	AAGTAGCTGG	TCAGCCTCCT	TCCTCCCACC	GCTCAAGCAA	TTGCCTTCAG
2280	GTTTTGACAT	TAGAGATGGG	TTGTTGTTTA	TAATTTTTTG	ACACCCTGGC	ATGCACCACC

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CAGAGGTGAC	ACTGAGACAC	CACTGGGCCT	GGAAATCAGG	GCATCAAGCC	AAAGAGGGTT	3240
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ACCTTCCTTC	TTTCCTATTT	CTCCATTTTC	TAATGGGAAT	GTCTATTATG	CCTGTTTCAC	3360
CATTGTACCT	TAGAAGCATG	TAACATTTCT	GGTTTCACAC	GTTCAAAGCT	GGAAAGGAAT	3420
TTTGTCTCTG	GATGAATCAC	ACATTGAGCC	TCACCCGTAA	CCTGATTTAG	ATGATTTTT	3480
AGATGACACT	TTGAACTTTA	GAATTGATGC	TAGAATGAGT	TAAGACTTTC	AGGGGGCTGT	3540
TGGGATGGAA	TAATTTTTTT	TTTTTTTTG	AGACGGAGTC	TAGCTCTGTC	GCCCAGGCTG	3600
GAGTGCAGTG	GCACCATCTT	GGCTCACTGC	AAGCTCTGCC	TCCCGGGTTT	ATGCCATTCT	3660
CATGTCTCAG	CCTCCAGAGT	AGCTGGGACT	ACAGGCGCCC	GCCACCACGC	CTGGCTAATT	3720
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TGACCTTCTG	ATCCGCCTGC	CTTGGCTTCC	CAAAGTGCTG	GGATTACAGG	TGTGAGCCAC	3840
CATGCCCGGC	TGGGATGGAA	TAAATTTATC	TTGTATGGGA	GAAGGACATA	CATTTTGGCA	3900

GGTCAAGGAC	AGAATGTTAT	GGACTAAACT	GTGTCCCCCA	AAATTCATTT	ATTAAAACCC	3960
TAAACCCCAG	TGTGACTGCA	TTTGGACATA	GAGCCTTTAG	GGGGTACATA	AAACTAAAGA	4020
TCACAGGATA	GGGCCCTAAT	CCCATTGGGG	CTGGTGTCCT	TACAGAAGAT	GAGACACTTA	4080
GAGCTCTCTC	TCCACGCAGG	CACCAAGGAA	ACACCATACA	AACACACAGT	GAGATGGCAG	4140
CCATCTGTTA	GCCAGGAACA	GATTCTCACC	ATAAACTATG	TTGGCACCTT	GATCTTAAAC	4200
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AAAAGATTCT	GTTGTTTAAG	CCATCCAGTC	TCTGGTATTT	TGTTATGGCA	GCCTGAGTAG	4320
GCTAAGACAA	TGAAGGATGT	GGTAAAACTT	TACGTCCCAA	CCACATACCA	AAGAGGCTGG	4380
AATTTAGCAT	GCTTTCTTCT	TTCAACTGTA	GGCAATGTGC	ACAAGTTCTA	AATCCTAAGA	4440
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CATCCAATGA	AGTTCTGACA	TTTCTTCAAC	ATGAGTACAG	TAATTCAATG	CCAGAGAATT	4560
CATTTTATTT	TGAAATCTAC	ATGCCATATT	CCAATTTCTG	TTGAAGATGC	AATGGTTATA	4620
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CAGTATTAAT	ACATTGTCAA	GATTTACCTC	TTCGTGTAGA	TTCCCTGGGG	AAAATTACCT	7080
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GAGTATCCAT	ACTAAACTCT	TTTCTACTGA	AAATAATGTG	CAAACATAAC	ATCCTATTCC	16380
TAGACAGTTT	GTAGTTTTTT	TCTCCCATTT	CTATTTTATA	AATCATCTTT	TTAAAATACT	16440
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GTAATAATTG	CCCTACTCAT	AAAGATGGGG	TGAAGATTAA	ATGAAATAGC	ACCTATAGAA	16620
CACTAGTTCC	AGACGTGGTA	TCATGCTAGT	AAAATGGCTG	CACAGCACTG	CTCAATGATG	16680
ACAAAAAGTG	AAGCTTCTGG	AGACAGACTC	CAAGTTTGAC	TCCCAGATCA	CCACATATAA	16740
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CTCTCTAGGC	ATATTCCTAA	GGACTTAAAG	AATGATAACT	ATCATTTCTC	TTAAATCTTC	17160
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GGGACTACAG	GTGCGCATGA	CTGTGACCAG	CTAATTTTTG	TATTTTTTTA	GAGACGGGTT	42180
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AAAACACCTC	TCCCCAGCTC	CAGCAACTGC	TGCAGCTGTG	CAGGGCAGTC	CCTCTCCAGG	42660
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GCCCTGGGTT	CTGCTGCTCT	CCAATCCAGT	GTGTCAGGGC	AGAATTCAAG	GTGGTCCTGC	42780

CCATCATACC	CGTACTTCCA	GTAGCCCTCG	GTACTGTTGT	CTTCTTGCAT	TTCACAGCCC	42840
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GGAATGCTGC	ACAGAGAGGC	CAAGAAGAAT	CTAGACAGAC	AGGCCTTGCT	GGGTTTCCCT	66660
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٠	TTGTTGGAGT	GCATTGGACA	TGGTAAAGTT	AAATATCGAT	TCATGAAACC	ATCGTTTGAG	217320
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CAAGTTAAAA	ATATTCAAAG	GAATGCCTAA	AGTTTTCAAA	ATTCTTTTAC	ATGTTGTACA	229440
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CGGAGTGCTG	AGAGCCTTCT	TAGTGTGTAT	TCAGTGTTTT	AAGAGAGCTT	GTGGATGAAT	232260
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CATTGGGGCC	AAATCTTAGA	TCATGTAAAT	TTTCTTCTAT	ATTTTATTCT	AAAAGCTTGT	235920
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TGTGTGTGT TGTGTGTG	237326
(2) INFORMATION FOR SEQ ID NO:3:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
CGTTAGAACG CGGCTACAAT	20
(2) INFORMATION FOR SEQ ID NO:4:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
•	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GCCGATTCAT TAATGCAGGT	20
(2) INFORMATION FOR SEQ ID NO:5:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

CTGTAAAACG ACGGCCAGTC	20
(2) INFORMATION FOR SEQ ID NO:6:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
GCAGGAAACA GCTATGACC	19
(2) INFORMATION FOR SEQ ID NO:7:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
GCATCAGCGA TTAACTTCTA C	21
(2) INFORMATION FOR SEQ ID NO:8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8: TTGCATTGTG GTGAAATCAG GG	22
(2) INFORMATION FOR SEQ ID NO:9:	
(i) SEQUENCE CHARACTERISTICS:	

(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
CTG	AGTAATT GTTTAAGGTG C	21
(2)	INFORMATION FOR SEQ ID NO:10:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
CTG	AGTAATT GTTTAAGGTG T	21
(2)	INFORMATION FOR SEQ ID NO:11:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
AGAZ	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	20
(2)	<pre>(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear</pre>	

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
GAA:	TGTGACC GTCCCATGAG	20
(2)	INFORMATION FOR SEQ ID NO:13:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
CAA	CTGAATA TGCAGAAAA AGTACACC	28
(2)	INFORMATION FOR SEQ ID NO:14:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
AGT	AGCTGGG ACTCACGGTG T	21
(2)	INFORMATION FOR SEQ ID NO:15:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

AGTAGCTGGG ACTCACGGTG C	21
(2) INFORMATION FOR SEQ ID NO:16:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
GCGCCACCAC TCCCAGCTCA T	21
(2) INFORMATION FOR SEQ ID NO:17:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17: TATATATATA TATA	14
(2) INFORMATION FOR SEQ ID NO:18:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
AAAAAAAAA AAAA	14
(2) INFORMATION FOR SEQ ID NO:19:	
(i) SEQUENCE CHARACTERISTICS:	

	(A) LENGTH: 10 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
AGGA	TGGTCT	10
(2)	INFORMATION FOR SEQ ID NO:20:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
TTGT	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	12
(2)	<pre>INFORMATION FOR SEQ ID NO:21: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
CTTT	TTTTTT TTTT	14
(2)	INFORMATION FOR SEQ ID NO:22:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

(XI) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
AATAATAATA AT	12
(2) INFORMATION FOR SEQ ID NO:23:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
TTTTTTTTT T	11
(2) INFORMATION FOR SEQ ID NO:24:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
ATATATATA ATATATATAT	20
AIAIAIAIA AIAIAIAIAI	20